Reviewer’s report

Title: Microorganisms involved in Deep Neck Infection (DNIs) in Greece: detection, identification and susceptibility to antimicrobials

Version: 0 Date: 23 Jul 2019

Reviewer: Jiun-Ling Wang

Reviewer's report:

P9 Line 39 "sequencing analysis failed to identify the microorganisms involved, given that a polymicrobial genetic pattern was obtained". Would you add some sentence about this in the discussion part and give as a limitation.

Table 2: Fusobacterium necrophorim and Actinomyces israelii were the predominant pathogen not identified by conventional culture method. Would you discuss more this about the reason? Is this related to anaerobic bacteria killed by antibiotic? or related to culture method? And what is the clinical implication after getting 16S rRNA PCR result? Is any underlying disease related to these two pathogen? Did this change the empirical antibiotic choice?

Are the methods appropriate and well described?
If not, please specify what is required in your comments to the authors.

Yes

Does the work include the necessary controls?
If not, please specify which controls are required in your comments to the authors.

Yes

Are the conclusions drawn adequately supported by the data shown?
If not, please explain in your comments to the authors.

Yes

Are you able to assess any statistics in the manuscript or would you recommend an additional statistical review?
If an additional statistical review is recommended, please specify what aspects require further assessment in your comments to the editors.

Not relevant to this manuscript

Quality of written English
Please indicate the quality of language in the manuscript:
Acceptable

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